1600

CRF Errors Corrected by the STIC Systems Branch

RECEIVED 7/15/03

Serial Number: 09/688,286D

CRF Processing Date: 7/15/03
Edited by: MT
Verified by: JUL 16 2003 (STIC staff)

TECH CENTER 1600

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data". ENTERED
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

1600

RAW SEQUENCE LISTING DATE: 07/15/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 07:51:52

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07152003\I688286D.raw**

See page 6.

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      3 <110> APPLICANT: Willson, Tracey
      4       Nicola , Nicos
      5       Hilton, Douglas
      6       Metcalf, Donald
      7       Zhang , Jian
      9 <120> TITLE OF INVENTION: A novel haemopoietin receptor and genetic sequences encoding
same
     11 <130> FILE REFERENCE: 23199-215
     13 <140> CURRENT APPLICATION NUMBER: US 09/688,286D
C--> 14 <141> CURRENT FILING DATE: 2003-07-10
     16 <150> PRIOR APPLICATION NUMBER: AU PN6135
     17 <151> PRIOR FILING DATE: 1995-10-23
     19 <150> PRIOR APPLICATION NUMBER: AU PN7276
     20 <151> PRIOR FILING DATE: 1995-12-22
     22 <150> PRIOR APPLICATION NUMBER: AU PP2208
     23 <151> PRIOR FILING DATE: 1996-09-09
     25 <160> NUMBER OF SEQ ID NOS: 12
     27 <170> SOFTWARE: PatentIn version 3.1
     29 <210> SEQ ID NO: 1
     30 <211> LENGTH: 1680
     31 <212> TYPE: DNA
     32 <213> ORGANISM: Mus musculus
     34 <220> FEATURE:
     35 <221> NAME/KEY: CDS
     36 <222> LOCATION: (61)..(1332)
     37 <223> OTHER INFORMATION:
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     43 Met Ala Arg Pro Ala Leu Leu Gly Glu Leu Leu Val Leu Leu Leu Trp
     44 1               5                  10                  15
     46 acc gcc acc gtg ggc caa gtt gcc gcg gcc aca gaa gtt cag cca cct      156
     47 Thr Ala Thr Val Gly Gln Val Ala Ala Ala Thr Glu Val Gln Pro Pro
     48             20                  25                  30
     50 gtg acg aat ttg agc gtc tct gtc gaa aat ctc tgc acg ata ata tgg      204
     51 Val Thr Asn Leu Ser Val Ser Val Glu Asn Leu Cys Thr Ile Ile Trp
     52         35                  40                  45
     54 acg tgg agt cct cct gaa gga gcc agt cca aat tgc act ctc aga tat      252
     55 Thr Trp Ser Pro Pro Glu Gly Ala Ser Pro Asn Cys Thr Leu Arg Tyr
     56     50                  55                  60
     58 ttt agt cac ttt gat gac caa cag gat aag aaa att gct cca gaa act      300
     59 Phe Ser His Phe Asp Asp Gln Gln Asp Lys Lys Ile Ala Pro Glu Thr
     60 65                  70                  75                  80
     62 cat cgt aaa gag gaa tta ccc ctg gat gag aaa atc tgt ctg cag gtg      348
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RAW SEQUENCE LISTING DATE: 07/15/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 07:51:52

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07152003\I688286D.raw**

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 63 His Arg Lys Glu Glu Leu Pro Leu Asp Glu Lys Ile Cys Leu Gln Val
 64                 85                  90                  95
 66 ggc tct cag tgt agt gcc aat gaa agt gag aag cct agc cct ttg gtg      396
 67 Gly Ser Gln Cys Ser Ala Asn Glu Ser Glu Lys Pro Ser Pro Leu Val
 68             100                 105                 110
 70 aaa aag tgc atc tca ccc cct gaa ggt gat cct gag tcc gct gtg act      444
 71 Lys Lys Cys Ile Ser Pro Pro Glu Gly Asp Pro Glu Ser Ala Val Thr
 72         115                 120                 125
 74 gag ctc aag tgc att tgg cat aac ctg agc tat atg aag tgt tcc tgg      492
 75 Glu Leu Lys Cys Ile Trp His Asn Leu Ser Tyr Met Lys Cys Ser Trp
 76     130                 135                 140
 78 ctc cct gga agg aat aca agc cct gac aca cac tat act ctg tac tat      540
 79 Leu Pro Gly Arg Asn Thr Ser Pro Asp Thr His Tyr Thr Leu Tyr Tyr
 80 145                 150                 155                 160
 82 tgg tac agc agc ctg gag aaa agt cgt caa tgt gaa aac atc tat aga      588
 83 Trp Tyr Ser Ser Leu Glu Lys Ser Arg Gln Cys Glu Asn Ile Tyr Arg
 84                 165                 170                 175
 86 gaa ggt caa cac att gct tgt tcc ttt aaa ttg act aaa gtg gaa cct      636
 87 Glu Gly Gln His Ile Ala Cys Ser Phe Lys Leu Thr Lys Val Glu Pro
 88             180                 185                 190
 90 agt ttt gaa cat cag aac gtt caa ata atg gtc aag gat aat gct ggg      684
 91 Ser Phe Glu His Gln Asn Val Gln Ile Met Val Lys Asp Asn Ala Gly
 92         195                 200                 205
 94 aaa att agg cca tcc tgc aaa ata gtg tct tta act tcc tat gtg aaa      732
 95 Lys Ile Arg Pro Ser Cys Lys Ile Val Ser Leu Thr Ser Tyr Val Lys
 96     210                 215                 220
 98 cct gat cct cca cat att aaa cat ctt ctc ctc aaa aat ggt gcc tta      780
 99 Pro Asp Pro Pro His Ile Lys His Leu Leu Leu Lys Asn Gly Ala Leu
100 225                 230                 235                 240
102 tta gtg cag tgg aag aat cca caa aat ttt aga agc aga tgc tta act      828
103 Leu Val Gln Trp Lys Asn Pro Gln Asn Phe Arg Ser Arg Cys Leu Thr
104                 245                 250                 255
106 tat gaa gtg gag gtc aat aat act caa acc gac cga cat aat att tta      876
107 Tyr Glu Val Glu Val Asn Asn Thr Gln Thr Asp Arg His Asn Ile Leu
108             260                 265                 270
110 gag gtt gaa gag gac aaa tgc cag aat tcc gaa tct gat aga aac atg      924
111 Glu Val Glu Glu Asp Lys Cys Gln Asn Ser Glu Ser Asp Arg Asn Met
112         275                 280                 285
114 gag ggt aca agt tgt ttc caa ctc cct ggt gtt ctt gcc gac gct gtc      972
115 Glu Gly Thr Ser Cys Phe Gln Leu Pro Gly Val Leu Ala Asp Ala Val
116     290                 295                 300
118 tac aca gtc aga gta aga gtc aaa aca aac aag tta tgc ttt gat gac     1020
119 Tyr Thr Val Arg Val Arg Val Lys Thr Asn Lys Leu Cys Phe Asp Asp
120 305                 310                 315                 320
122 aac aaa ctg tgg agt gat tgg agt gaa gca cag agt ata ggt aag gag     1068
123 Asn Lys Leu Trp Ser Asp Trp Ser Glu Ala Gln Ser Ile Gly Lys Glu
124                 325                 330                 335
126 caa aac tcc acc ttc tac acc acc atg tta ctc acc att cca gtc ttt     1116
127 Gln Asn Ser Thr Phe Tyr Thr Thr Met Leu Leu Thr Ile Pro Val Phe
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RAW SEQUENCE LISTING DATE: 07/15/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 07:51:52

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07152003\I688286D.raw**

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128                 340                 345                 350
130 gtc gca gtg gca gtc ata atc ctc ctt ttt tac ctg aaa agg ctt aag     1164
131 Val Ala Val Ala Val Ile Ile Leu Leu Phe Tyr Leu Lys Arg Leu Lys
132         355                 360                 365
134 atc att ata ttt cct cca att cct gat cct ggc aag att ttt aaa gaa     1212
135 Ile Ile Ile Phe Pro Pro Ile Pro Asp Pro Gly Lys Ile Phe Lys Glu
136     370                 375                 380
138 atg ttt gga gac cag aat gat gat acc ctg cac tgg aag aag tat gac     1260
139 Met Phe Gly Asp Gln Asn Asp Asp Thr Leu His Trp Lys Lys Tyr Asp
140 385                 390                 395                 400
142 atc tat gag aaa caa tcc aaa gaa gaa acg gat tct gta gtg ctg ata     1308
143 Ile Tyr Glu Lys Gln Ser Lys Glu Glu Thr Asp Ser Val Val Leu Ile
144                 405                 410                 415
146 gaa aac ctg aag aaa gca gct cct tgatggggag aagtgatttc tttcttgcct    1362
147 Glu Asn Leu Lys Lys Ala Ala Pro
148             420
150 tcaatgtgac cctgtgaaga tttattgcat tctccatttg ttatctgggg gacttgttaa   1422
152 atagaaactg aaactactct tgaaaaacag gcagctccta agagccacag gtcttgatgt   1482
154 gacttttgca ttgaaaaccc aaacccaaag gagctccttc caagaaaagc aagagttctt   1542
156 ctcgttcctt gttccaatcc ctaaaagcag atgttttgcc aaatccccaa actagaggac   1602
158 aaagacaagg ggacaatgac catcaattca tctaatcagg aattgtgatg gcttcctaag   1662
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163 <210> SEQ ID NO: 2
164 <211> LENGTH: 424
165 <212> TYPE: PRT
166 <213> ORGANISM: Mus musculus
168 <400> SEQUENCE: 2
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175             20                  25                  30
178 Val Thr Asn Leu Ser Val Ser Val Glu Asn Leu Cys Thr Ile Ile Trp
179         35                  40                  45
182 Thr Trp Ser Pro Pro Glu Gly Ala Ser Pro Asn Cys Thr Leu Arg Tyr
183     50                  55                  60
186 Phe Ser His Phe Asp Asp Gln Gln Asp Lys Lys Ile Ala Pro Glu Thr
187 65                  70                  75                  80
190 His Arg Lys Glu Glu Leu Pro Leu Asp Glu Lys Ile Cys Leu Gln Val
191                 85                  90                  95
194 Gly Ser Gln Cys Ser Ala Asn Glu Ser Glu Lys Pro Ser Pro Leu Val
195             100                 105                 110
198 Lys Lys Cys Ile Ser Pro Pro Glu Gly Asp Pro Glu Ser Ala Val Thr
199         115                 120                 125
202 Glu Leu Lys Cys Ile Trp His Asn Leu Ser Tyr Met Lys Cys Ser Trp
203     130                 135                 140
206 Leu Pro Gly Arg Asn Thr Ser Pro Asp Thr His Tyr Thr Leu Tyr Tyr
207 145                 150                 155                 160
210 Trp Tyr Ser Ser Leu Glu Lys Ser Arg Gln Cys Glu Asn Ile Tyr Arg
211                 165                 170                 175
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RAW SEQUENCE LISTING DATE: 07/15/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 07:51:52

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07152003\I688286D.raw**

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215             180                 185                 190
218 Ser Phe Glu His Gln Asn Val Gln Ile Met Val Lys Asp Asn Ala Gly
219         195                 200                 205
222 Lys Ile Arg Pro Ser Cys Lys Ile Val Ser Leu Thr Ser Tyr Val Lys
223     210                 215                 220
226 Pro Asp Pro Pro His Ile Lys His Leu Leu Leu Lys Asn Gly Ala Leu
227 225                 230                 235                 240
230 Leu Val Gln Trp Lys Asn Pro Gln Asn Phe Arg Ser Arg Cys Leu Thr
231                 245                 250                 255
234 Tyr Glu Val Glu Val Asn Asn Thr Gln Thr Asp Arg His Asn Ile Leu
235             260                 265                 270
238 Glu Val Glu Glu Asp Lys Cys Gln Asn Ser Glu Ser Asp Arg Asn Met
239         275                 280                 285
242 Glu Gly Thr Ser Cys Phe Gln Leu Pro Gly Val Leu Ala Asp Ala Val
243     290                 295                 300
246 Tyr Thr Val Arg Val Arg Val Lys Thr Asn Lys Leu Cys Phe Asp Asp
247 305                 310                 315                 320
250 Asn Lys Leu Trp Ser Asp Trp Ser Glu Ala Gln Ser Ile Gly Lys Glu
251                 325                 330                 335
254 Gln Asn Ser Thr Phe Tyr Thr Thr Met Leu Leu Thr Ile Pro Val Phe
255             340                 345                 350
258 Val Ala Val Ala Val Ile Ile Leu Leu Phe Tyr Leu Lys Arg Leu Lys
259         355                 360                 365
262 Ile Ile Ile Phe Pro Pro Ile Pro Asp Pro Gly Lys Ile Phe Lys Glu
263     370                 375                 380
266 Met Phe Gly Asp Gln Asn Asp Asp Thr Leu His Trp Lys Lys Tyr Asp
267 385                 390                 395                 400
270 Ile Tyr Glu Lys Gln Ser Lys Glu Glu Thr Asp Ser Val Val Leu Ile
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275             420
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279 <211> LENGTH: 1383
280 <212> TYPE: DNA
281 <213> ORGANISM: human
283 <220> FEATURE:
284 <221> NAME/KEY: CDS
285 <222> LOCATION: (61)..(1338)
286 <223> OTHER INFORMATION:
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291 atg gag tgg ccg gcg cgg ctc tgc ggg ctg tgg gcg ctg ctg ctc tgc      108
292 Met Glu Trp Pro Ala Arg Leu Cys Gly Leu Trp Ala Leu Leu Leu Cys
293 1               5                   10                  15
295 gcc ggc ggc ggg ggc ggg ggc ggg ggc gcg cct acg gaa act cag cca      156
296 Ala Gly Gly Gly Gly Gly Gly Gly Gly Ala Pro Thr Glu Thr Gln Pro
297             20                  25                  30
299 cct gtg aca aat ttg agt gtc tct gtt gaa aac ctc tgc aca gta ata      204
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RAW SEQUENCE LISTING DATE: 07/15/2003
PATENT APPLICATION: US/09/688,286D TIME: 07:51:52

Input Set : A:\11373A.seqlist.txt
Output Set: N:\CRF4\07152003\I688286D.raw

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300 Pro Val Thr Asn Leu Ser Val Ser Val Glu Asn Leu Cys Thr Val Ile
301         35                  40                  45
303 tgg aca tgg aat cca ccc gag gga gcc agc tca aat tgt agt cta tgg      252
304 Trp Thr Trp Asn Pro Pro Glu Gly Ala Ser Ser Asn Cys Ser Leu Trp
305     50                  55                  60
307 tat ttt agt cat ttt ggc gac aaa caa gat aag aaa ata gct ccg gaa      300
308 Tyr Phe Ser His Phe Gly Asp Lys Gln Asp Lys Lys Ile Ala Pro Glu
309 65                  70                  75                  80
311 act cgt cgt tca ata gaa gta ccc ctg aat gag agg att tgt ctg caa      348
312 Thr Arg Arg Ser Ile Glu Val Pro Leu Asn Glu Arg Ile Cys Leu Gln
313                 85                  90                  95
315 gtg ggg tcc cag tgt agc acc aat gag agt gag aag cct agc att ttg      396
316 Val Gly Ser Gln Cys Ser Thr Asn Glu Ser Glu Lys Pro Ser Ile Leu
317             100                 105                 110
319 gtt gaa aaa tgc atc tca ccc cca gaa ggt gat cct gag tct gct gtg      444
320 Val Glu Lys Cys Ile Ser Pro Pro Glu Gly Asp Pro Glu Ser Ala Val
321         115                 120                 125
323 act gaa ctt caa tgc att tgg cac aac ctg agc tac atg aag tgt tct      492
324 Thr Glu Leu Gln Cys Ile Trp His Asn Leu Ser Tyr Met Lys Cys Ser
325     130                 135                 140
327 tgg ctc cct gga agg aat acc agt ccc gac act aac tat act ctc tac      540
328 Trp Leu Pro Gly Arg Asn Thr Ser Pro Asp Thr Asn Tyr Thr Leu Tyr
329 145                 150                 155                 160
331 tat tgg cac aga agc ctg gaa aaa att cat caa tgt gaa aac atc ttt      588
332 Tyr Trp His Arg Ser Leu Glu Lys Ile His Gln Cys Glu Asn Ile Phe
333                 165                 170                 175
335 aga gaa ggc caa tac ttt ggt tgt tcc ttt gat ctg acc aaa gtg aag      636
336 Arg Glu Gly Gln Tyr Phe Gly Cys Ser Phe Asp Leu Thr Lys Val Lys
337             180                 185                 190
339 gat tcc agt ttt gaa caa cac agt gtc caa ata atg gtc aag gat aat      684
340 Asp Ser Ser Phe Glu Gln His Ser Val Gln Ile Met Val Lys Asp Asn
341         195                 200                 205
343 gca gga aaa att aaa cca tcc ttc aat ata gtg cct tta act tcc cgt      732
344 Ala Gly Lys Ile Lys Pro Ser Phe Asn Ile Val Pro Leu Thr Ser Arg
345     210                 215                 220
347 gtg aaa cct gat cct cca cat att aaa aac ctc tcc ttc cac aat gat      780
348 Val Lys Pro Asp Pro Pro His Ile Lys Asn Leu Ser Phe His Asn Asp
349 225                 230                 235                 240
351 gac cta tat gtg caa tgg gag aat cca cag aat ttt att agc aga tgc      828
352 Asp Leu Tyr Val Gln Trp Glu Asn Pro Gln Asn Phe Ile Ser Arg Cys
353                 245                 250                 255
355 cta ttt tat gaa gta gaa gtc aat aac agc caa act gag aca cat aat      876
356 Leu Phe Tyr Glu Val Glu Val Asn Asn Ser Gln Thr Glu Thr His Asn
357             260                 265                 270
359 gtt ttc tac gtc caa gag gct aaa tgt gag aat cca gaa ttt gag aga      924
360 Val Phe Tyr Val Gln Glu Ala Lys Cys Glu Asn Pro Glu Phe Glu Arg
361         275                 280                 285
363 aat gtg gag aat aca tct tgt ttc atg gtc cct ggt gtt ctt cct gat      972
364 Asn Val Glu Asn Thr Ser Cys Phe Met Val Pro Gly Val Leu Pro Asp
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/15/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 07:51:53

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07152003\I688286D.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; Xaa Pos. 3
Seq#:10; Xaa Pos. 24
Seq#:11; Xaa Pos. 24

VERIFICATION SUMMARY DATE: 07/15/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 07:51:53

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07152003\I688286D.raw**

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:39 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:37
L:288 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:286
L:589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:613 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:16
L:637 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:16

1600

RAW SEQUENCE LISTING DATE: 07/10/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 11:05:02

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07102003\I688286D.raw**

3 <110> APPLICANT: Willson, Tracey
4 Nicola , Nicos
5 Hilton, Douglas
6 Metcalf, Donald
7 Zhang , Jian
9 <120> TITLE OF INVENTION: A novel haemopoietin receptor and genetic sequences encoding same
11 <130> FILE REFERENCE: 23199-215
13 <140> CURRENT APPLICATION NUMBER: US 09/688,286D
C--> **14 <141> CURRENT FILING DATE: 2000-10-13**
16 <150> PRIOR APPLICATION NUMBER: AU PN6135
17 <151> PRIOR FILING DATE: 1995-10-23
19 <150> PRIOR APPLICATION NUMBER: AU PN7276
20 <151> PRIOR FILING DATE: 1995-12-22
22 <150> PRIOR APPLICATION NUMBER: AU PP2208
23 <151> PRIOR FILING DATE: 1996-09-09
25 <160> NUMBER OF SEQ ID NOS: 12
27 <170> SOFTWARE: PatentIn version 3.1

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

641 <210> SEQ ID NO: 12
642 <211> LENGTH: 5
643 <212> TYPE: PRT
644 <213> ORGANISM: unknown
646 <220> FEATURE:
647 <223> OTHER INFORMATION: peptide motif found in many members of the haemopoietin receptor
648 family
650 <400> SEQUENCE: 12
652 Trp Ser Asp Trp Ser
653 1 5
E--> 662 (- 11 -) Delete

VERIFICATION SUMMARY DATE: 07/10/2003
PATENT APPLICATION: **US/09/688,286D** TIME: 11:05:03

Input Set : **A:\11373A.seqlist.txt**
Output Set: **N:\CRF4\07102003\I688286D.raw**

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:39 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:37
L:288 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:286
L:589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:613 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:16
L:637 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:16
L:662 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:12